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RAW SEQUENCE LISTING

DATE: 05/07/2002

PATENT APPLICATION: US/09/943,115A

TIME: 14:51:46

Input Set : D:\52459-20021.txt

Output Set: N:\CRF3\05072002\I943115A.raw

ENTERED

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4 <110> APPLICANT: SEQUENOM, Inc.
5     Risinger, Carl
6     Andersson, Maria
7     Lewander, Tommy
8     Olaisson, Erik
10 <120> TITLE OF INVENTION: DETECTION OF CYP3A4 AND CYP2C9
11     POLYMORPHISMS
13 <130> FILE REFERENCE: 52459-20021.00
15 <140> CURRENT APPLICATION NUMBER: US 09/943,115A
16 <141> CURRENT FILING DATE: 2001-08-30
18 <150> PRIOR APPLICATION NUMBER: UK 0021286.0
19 <151> PRIOR FILING DATE: 2000-08-30
21 <160> NUMBER OF SEQ ID NOS: 73
23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 1345
27 <212> TYPE: DNA
28 <213> ORGANISM: Homo sapiens
30 <400> SEQUENCE: 1
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32 gggcagctgt tctcttctct cctttctctc ctgtttccag acatgcagta ttccagaga      120
33 gaaggggcca ctctttggca aagaacctgt ctaacttgct atctatggca ggacctttga      180
34 aggggttcaca ggaagcagca caaattgata ctattccacc aagccatcag ctccatctca      240
35 tccatgccct gtctctcctt taggggtccc cttgccaaaca gaatcacaga ggaccagcct      300
36 gaaagtgcag agacagcagc tgaggcacag ccaagagctc tggctgtatt aatgacctaa      360
37 gaagtcacca gaaagtcaga aggatgcata gcagaggccc agcaatctca gctaagtcaa      420
38 ctccaccagc ctttctagtt gcccaactgt tgtacagcac sctggtaggg accagagcca      480
39 tgacagggaa taagactaga ctatgccctt gaggagctca cctctgttca gggaaacagg      540
40 cgtggaaaca caatggtggt aaagaggaaa gaggacaata ggattgcatg aaggggatgg      600
41 aaagtgccca ggggaggaaa tgggtacatc tgtgtgagga gtttgggtgag gaaagactct      660
42 aagagaaggc tctgtctgtc tgggtttgga aggatgtgta ggagtcttct agggggcaca      720
43 ggcacactcc aggcataagg aaagatctgt aggtgtggct tgttgggatg aatttcaagt      780
44 attttggaat gaggacagcc atagagacaa gggcargaga gaggcgattt aatagatttt      840
45 atgccaatgg ctccacttga gtttctgata agaaccaga acccttgga cccccagtaa      900
46 cattgattga gttgtttatg atacctcata gaatatgaac tcaaaggagg tcagtgagtg      960
47 gtgtgtgtgt gattctttgc caacttccaa ggtggagaag cctcttccaa ctgcaggcag      1020
48 agcacaggtg gccctgttac tggctgcagc tccagccctg cctccttctc tagcatataa      1080
49 acaatccaac agcctcactg aatcactgct tgcaggggca ggaaagctcc atgcacatag      1140
50 cccagcaaaag agcaacacag agctgaaaagg aagactcaga ggagagagat aagtaaggaa      1200
51 agtagtgatg gctctcatcc cagacttggc catggaaaacc tggcttctcc tggctgtcag      1260
52 cctggtgtct ctctatctgt gagtaactgt tcaggctcct cttctctgtt tcttggactt      1320
53 ggggtcgtaa tcaggcctct cttttt
55 <210> SEQ ID NO: 2

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56 <211> LENGTH: 19
57 <212> TYPE: DNA
58 <213> ORGANISM: Artificial Sequence
60 <220> FEATURE:
61 <223> OTHER INFORMATION: Oligonucleotide of CYP3A4 region
63 <400> SEQUENCE: 2
64 acaagggcaa gagagaggc 19
66 <210> SEQ ID NO: 3
67 <211> LENGTH: 19
68 <212> TYPE: DNA
69 <213> ORGANISM: Artificial Sequence
71 <220> FEATURE:
72 <223> OTHER INFORMATION: Oligonucleotide of CYP3A4 region
74 <400> SEQUENCE: 3
75 acaagggcag gagagaggc 19
77 <210> SEQ ID NO: 4
78 <211> LENGTH: 10
79 <212> TYPE: DNA
80 <213> ORGANISM: Artificial Sequence
82 <220> FEATURE:
83 <223> OTHER INFORMATION: Oligonucleotide of CYP3A4 region
85 <400> SEQUENCE: 4
86 agggcaagag 10
88 <210> SEQ ID NO: 5
89 <211> LENGTH: 10
90 <212> TYPE: DNA
91 <213> ORGANISM: Artificial Sequence
93 <220> FEATURE:
94 <223> OTHER INFORMATION: Oligonucleotide of CYP3A4 region
96 <400> SEQUENCE: 5
97 agggcaggag 10
99 <210> SEQ ID NO: 6
100 <211> LENGTH: 2438
101 <212> TYPE: DNA
102 <213> ORGANISM: Homo sapiens
104 <400> SEQUENCE: 6
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106 aggtcctaga aggagccgca gtcagcagg agagaggagg agctgagctg ggaccctac 120
107 ctcttgagga atgaaatgat tattataaag acagcaaccg agcttatttt acccaaaata 180
108 aggtagtata tttctgttag agtttagagt ttcatgagtc agggaccaag ttattgcttt 240
109 tctttgccct gtataaaggc ttctccaagg cctttgaact acctaagtac taaatgttat 300
110 aaaaccaaac tcttctgacc tctcaatcta gtcaactggg gctgtaatta ttaatgaaat 360
111 taatgtttat ttgaaaata atttactaga ctgaattacg aaatcctgaa tcattgtaca 420
112 ctatcagtaa atattggtgg acccaactga actgaatggt ttgcttgaaa tgaaaccttt 480
113 gagatgcagg gcttatgggt tctagtccca gctctagcac tagcagacag catgttcttg 540
114 gctaagatac tgaatcttca aggtcagct tcctcattcc ggaaatgggt caattttatt 600
115 gtaagcagag gtaattgaga gattcaaaag ggacatgagg tgtaacaatt ctctgtaaat 660
116 tgttagaatc cctgttaaaa atgaccagta aagctttgtg caactgtgtc ttgacataac 720
117 tttatttttc ttaataaaaag aaatggaaat aacctcacta gggaatttag aacaaatatg 780

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118 atgatatctt taaagaaaaat ggctttgcac aagtattgac attaatgac tagtaaagtg      840
119 tatctttcta gttgtattta gatcctcaac tcagtatgtc agtcctgtt aaggctctata      900
120 cattgtggtg gttctgtgct gtgggtccat ttagtgattt ccctacctcc catcttytat      960
121 tgcattccaca actgtggttc tgcocataat ttcttttgc tctgtgcat tattacatca     1020
122 tatctgaaaa tgagaaacca aaaacaatrg aaagcagcca tgtctggagg tgactggggg     1080
123 gtcgagaagc cctagtttct caaacccctta gcaccaaatt ttccctcag ttacactgag     1140
124 cgtttcactt ctgcagtgat ggaraaggga gatcccttat ttcttctcat gagcatctct     1200
125 ggtgctgttt cccttagaga caaataaggg gttctattta atgtgaagcc tgttttatga     1260
126 acagaataaa tgtggtgtat attcagaata actaatgttt ggaagttgtt ttatttttgc     1320
127 taaaaattgt tctcaaggca gctctggtgt aagagataat acaccacgat gggcatcaga     1380
128 agacctcagc tcaaatccca gttctgccag ctatgagctg tgtggcacca acaggtgtcc     1440
129 tgttctccca ggtctccct tttcccattt gaaaaataaa aaataacaat tctgccttc     1500
130 aggaattttt tttagggggg ttaatkgtaa aggtgtttat atctgctaag gtaatttact     1560
131 tgatatatgt ttggttattt aagatatatg agttatgtta gctatttcat gtttaggctg     1620
132 ctgtattttt agtaggctat attaaatatt tgaaaggatt wmattataaa gaacaaagtc     1680
133 tctaatctt tgatatagca ttgacatact ttttaatat acaaggcata gaatatggcc     1740
134 atttctgtta aatcatatat tcccaactgg ttattaatct aagaattcag aattttgagt     1800
135 aattgctttt gcatcagatt atttacttca gtgctctcaa ttatgatggt gcattagaac     1860
136 catctgggtt aacatttggt ttttattacc aatacctagg ctccaacca gtacagtga     1920
137 actggaatgt acagagtgga caatggaacg aaggagaaca agaccaaagg acattttatt     1980
138 tttatctgta tcagtgggtc aaagtccttt cagaaggagc atatagtgga cctaggtgat     2040
139 tggtaattt atccatcaaa gaggcacaca ccgaattagc atggagtgtt ataaaaggct     2100
140 tggagtgcaa gctcatggtt gtcttaacaa gaagagaagg cttcaatgga ttctcttggt     2160
141 gtcttctgtc tctgtctctc atgtttgctt ctcccttcac tctggagaca gagctctggg     2220
142 agaggaaaac tccctcctgg ccccaactct ctcccagtga ttggaaatat cctacagata     2280
143 ggtattaagg acatcagcaa atccttaacc aatgtaagta tgctccttca gtggcttgca     2340
144 aaaggtaagt aaattcacct gtatttttta aataaagtgt atccctagag gtacatgtta     2400
145 caagaggtaa tggtaaagta aaatactttg aaaggctt     2438

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147 <210> SEQ ID NO: 7

148 <211> LENGTH: 20

149 <212> TYPE: DNA

150 <213> ORGANISM: Artificial Sequence

152 <220> FEATURE:

153 <223> OTHER INFORMATION: Primer

155 <400> SEQUENCE: 7

156 ccagcctgaa agtgcagaga

20

158 <210> SEQ ID NO: 8

159 <211> LENGTH: 25

160 <212> TYPE: DNA

161 <213> ORGANISM: Artificial Sequence

163 <220> FEATURE:

164 <223> OTHER INFORMATION: Primer

166 <400> SEQUENCE: 8

167 tcttagagtc ttctctcacc aaact

25

169 <210> SEQ ID NO: 9

170 <211> LENGTH: 20

171 <212> TYPE: DNA

172 <213> ORGANISM: Artificial Sequence

174 <220> FEATURE:

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178 catgccctgt ctctccttta                                20
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182 <212> TYPE: DNA
183 <213> ORGANISM: Artificial Sequence
185 <220> FEATURE:
186 <223> OTHER INFORMATION: Primer
188 <400> SEQUENCE: 10
189 ccatcccctt catgcaatc                                19
191 <210> SEQ ID NO: 11
192 <211> LENGTH: 11
193 <212> TYPE: DNA
194 <213> ORGANISM: Artificial Sequence
196 <220> FEATURE:
197 <223> OTHER INFORMATION: Oligonucleotide representing the coding strand
198     complementary to the polymorphic site 461
200 <400> SEQUENCE: 11
201 agcaccctgg t                                        11
203 <210> SEQ ID NO: 12
204 <211> LENGTH: 11
205 <212> TYPE: DNA
206 <213> ORGANISM: Artificial Sequence
208 <220> FEATURE:
209 <223> OTHER INFORMATION: Oligonucleotide representing the coding strand
210     complementary to the polymorphic site 461
212 <400> SEQUENCE: 12
213 agcacgctgg t                                        11
215 <210> SEQ ID NO: 13
216 <211> LENGTH: 11
217 <212> TYPE: DNA
218 <213> ORGANISM: Artificial Sequence
220 <220> FEATURE:
221 <223> OTHER INFORMATION: Oligonucleotide representing the non-coding strand
222     complementary to the polymorphic site 461
224 <400> SEQUENCE: 13
225 accaggggtgc t                                        11
227 <210> SEQ ID NO: 14
228 <211> LENGTH: 11
229 <212> TYPE: DNA
230 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: Oligonucleotide representing the non-coding strand
234     complementary to the polymorphic site 461
236 <400> SEQUENCE: 14
237 accagcgtgc t                                        11
239 <210> SEQ ID NO: 15
240 <211> LENGTH: 11

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241 <212> TYPE: DNA
242 <213> ORGANISM: Artificial Sequence
244 <220> FEATURE:
245 <223> OTHER INFORMATION: Oligonucleotide of the novel polymorphic site 461
246     on the coding strand
248 <400> SEQUENCE: 15
249 gtgtgtacag c                                     11
251 <210> SEQ ID NO: 16
252 <211> LENGTH: 11
253 <212> TYPE: DNA
254 <213> ORGANISM: Artificial Sequence
256 <220> FEATURE:
257 <223> OTHER INFORMATION: Oligonucleotide of the novel polymorphic site 461
258     on the non-coding strand
260 <400> SEQUENCE: 16
261 gctgtacaca c                                     11
263 <210> SEQ ID NO: 17
264 <211> LENGTH: 11
265 <212> TYPE: DNA
266 <213> ORGANISM: Artificial Sequence
268 <220> FEATURE:
269 <223> OTHER INFORMATION: Oligonucleotide of the novel polymorphic site 461
270     on the non-coding strand
272 <400> SEQUENCE: 17
273 tggtccttac c                                     11
275 <210> SEQ ID NO: 18
276 <211> LENGTH: 11
277 <212> TYPE: DNA
278 <213> ORGANISM: Artificial Sequence
280 <220> FEATURE:
281 <223> OTHER INFORMATION: Oligonucleotide of the novel polymorphic site 461
282     on the coding strand
284 <400> SEQUENCE: 18
285 ggtagggacc a                                     11
287 <210> SEQ ID NO: 19
288 <211> LENGTH: 25
289 <212> TYPE: DNA
290 <213> ORGANISM: Artificial Sequence
292 <220> FEATURE:
293 <223> OTHER INFORMATION: Primer
295 <400> SEQUENCE: 19
296 cactagggaa tttagaaca ataatg                       25
298 <210> SEQ ID NO: 20
299 <211> LENGTH: 23
300 <212> TYPE: DNA
301 <213> ORGANISM: Artificial Sequence
303 <220> FEATURE:
304 <223> OTHER INFORMATION: Primer
306 <400> SEQUENCE: 20

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VERIFICATION SUMMARY

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